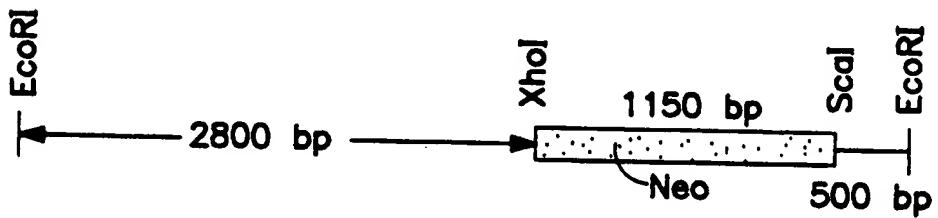
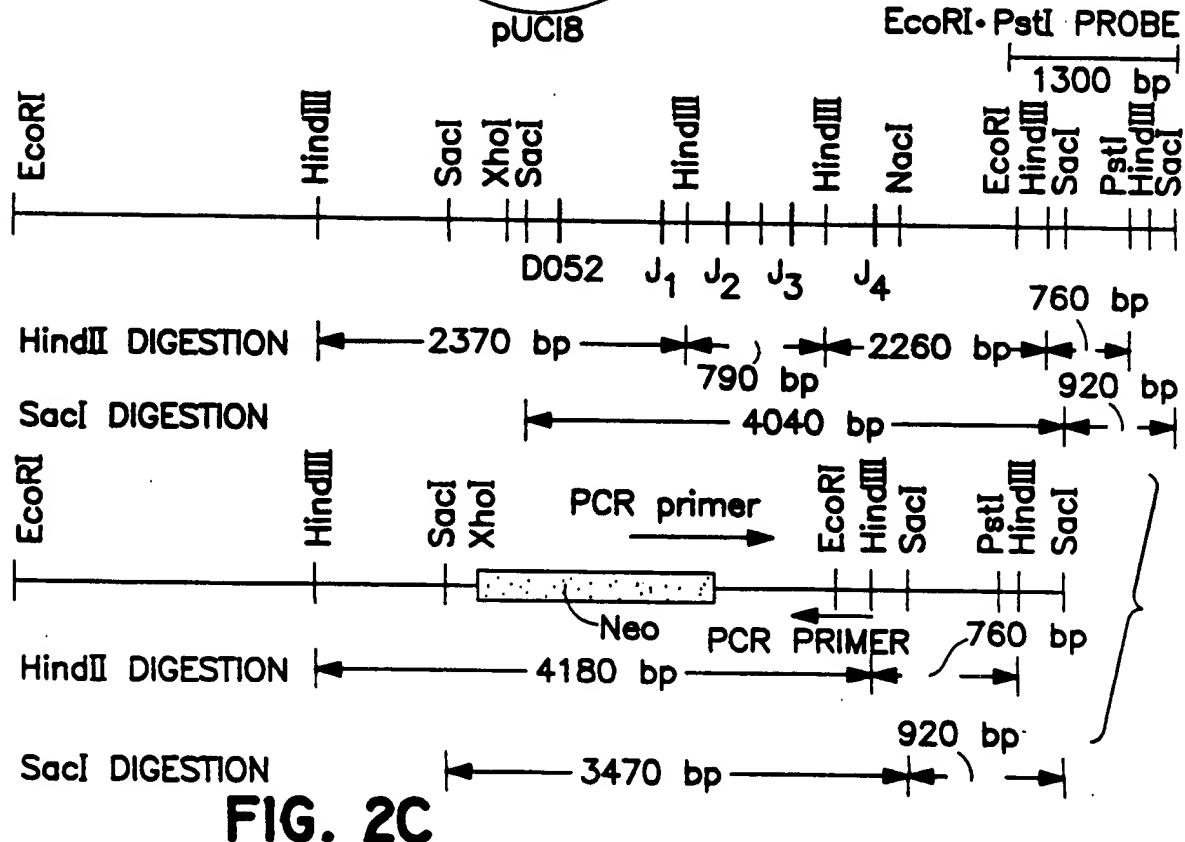
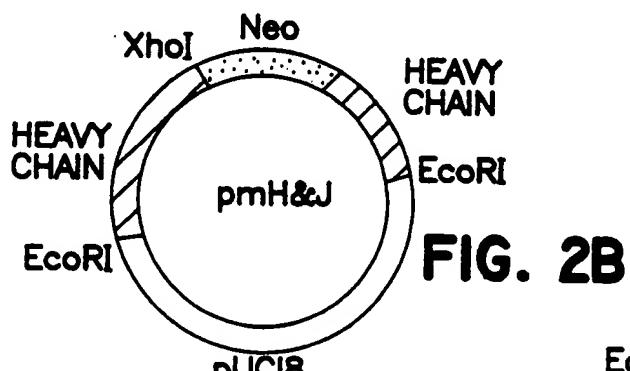
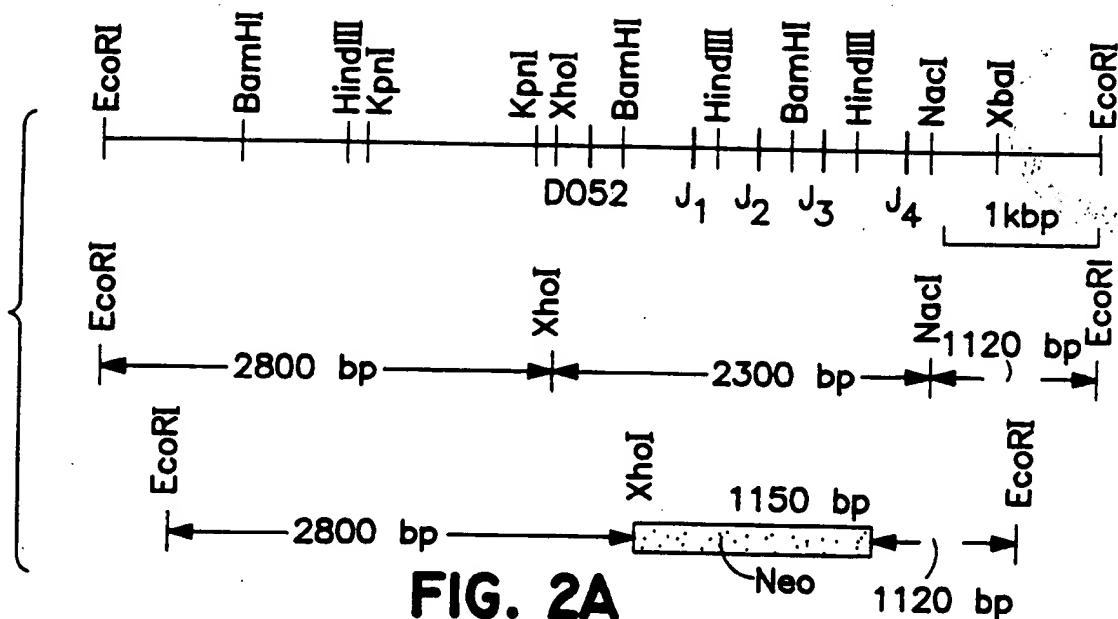
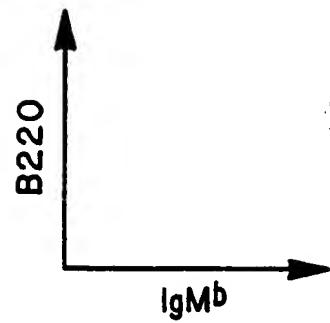
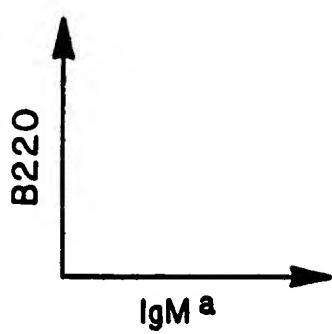
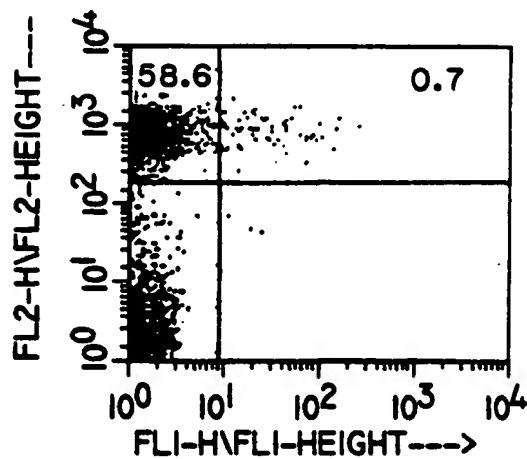
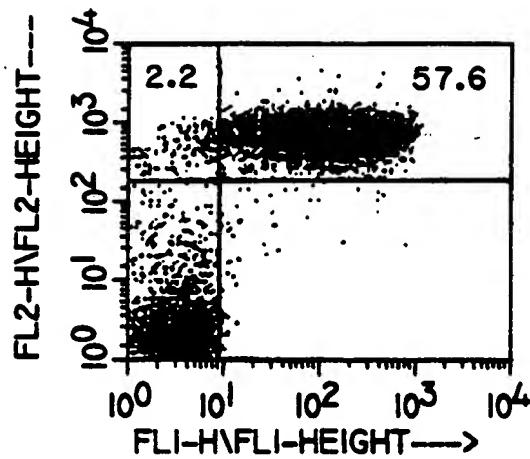
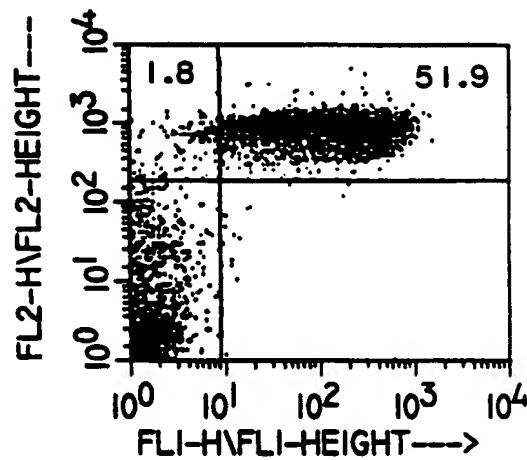
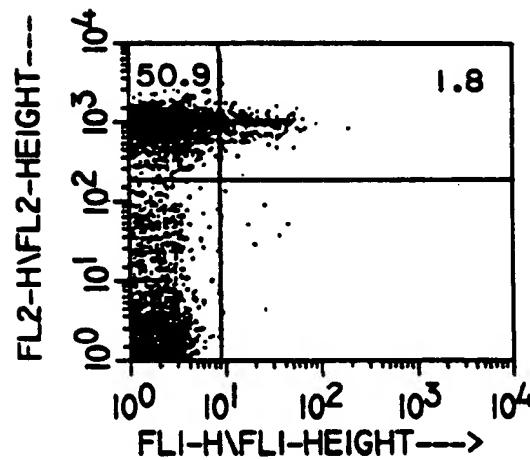
**FIG. IA****FIG. IB**



**a allotype****b allotype****FIG. 3-1**

a/b F1

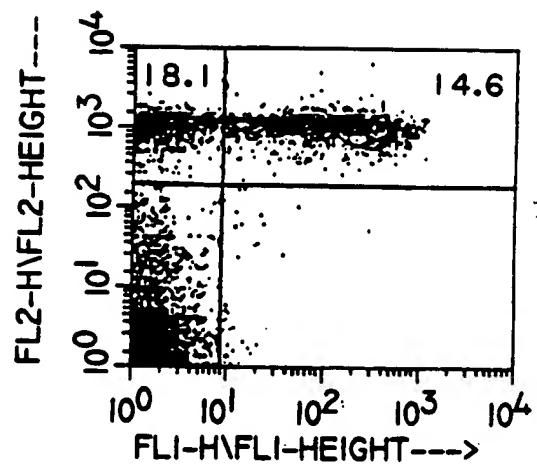
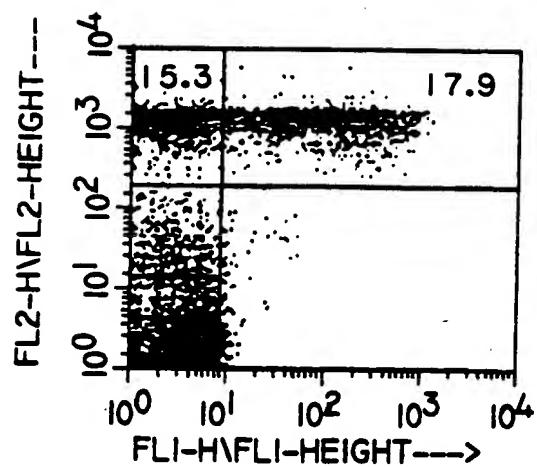
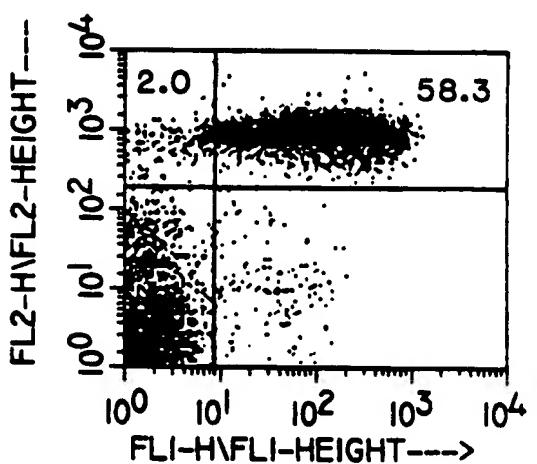
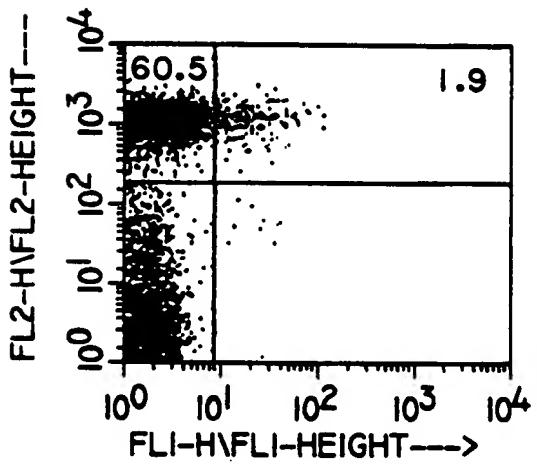
 $\Delta J_H/b$ F1

FIG. 3-2

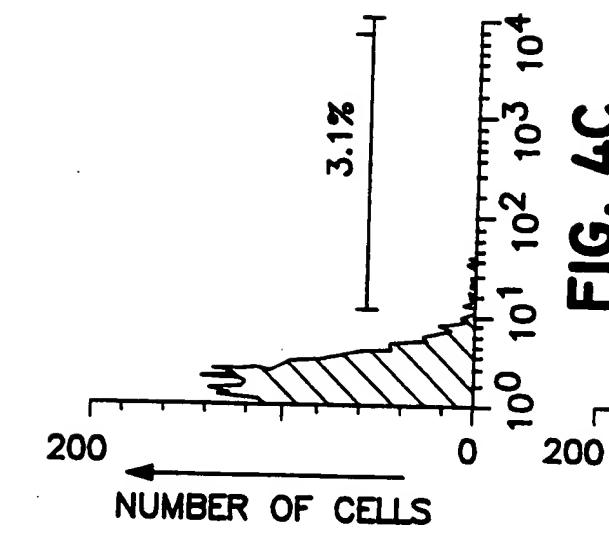


FIG. 4C

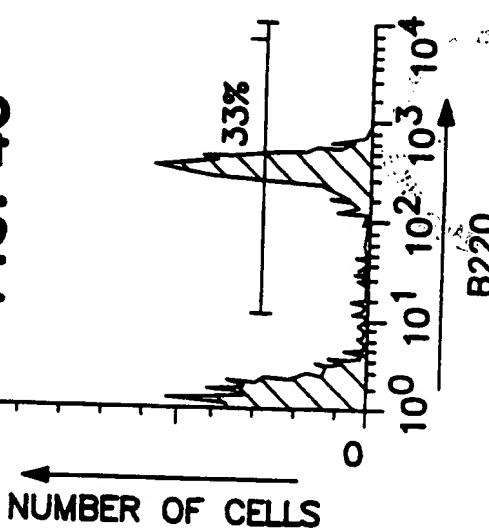


FIG. 4F

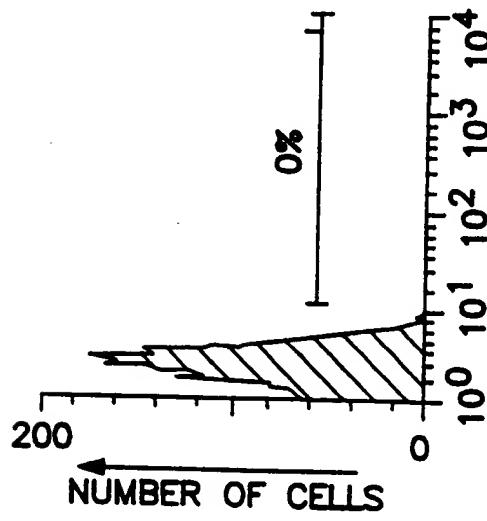


FIG. 4B

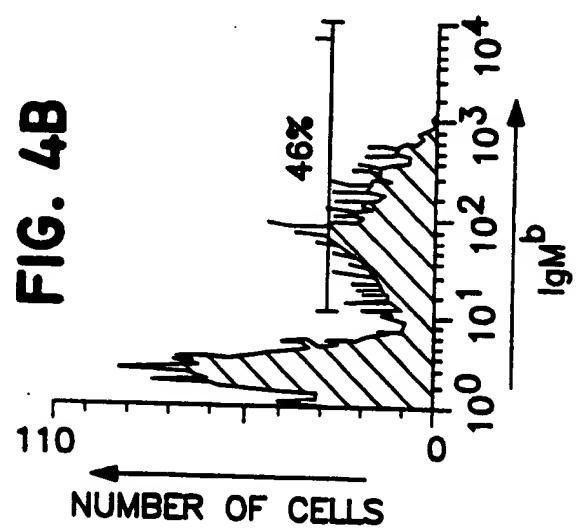


FIG. 4E

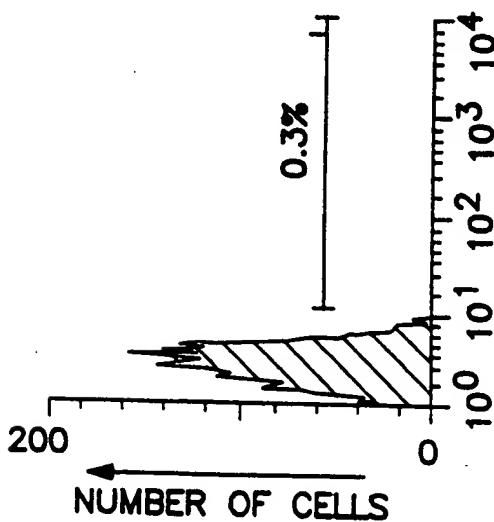


FIG. 4A

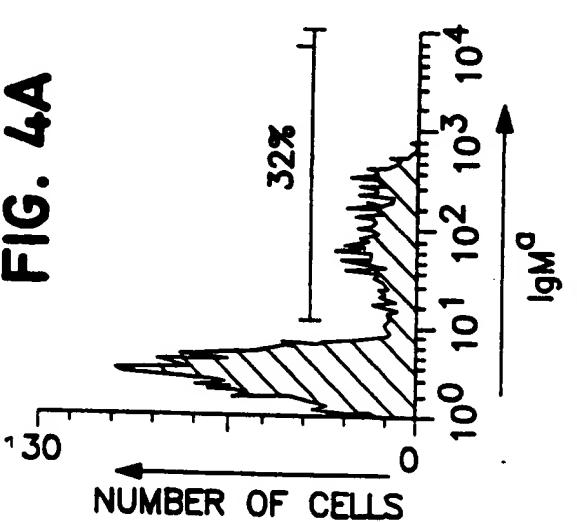


FIG. 4D

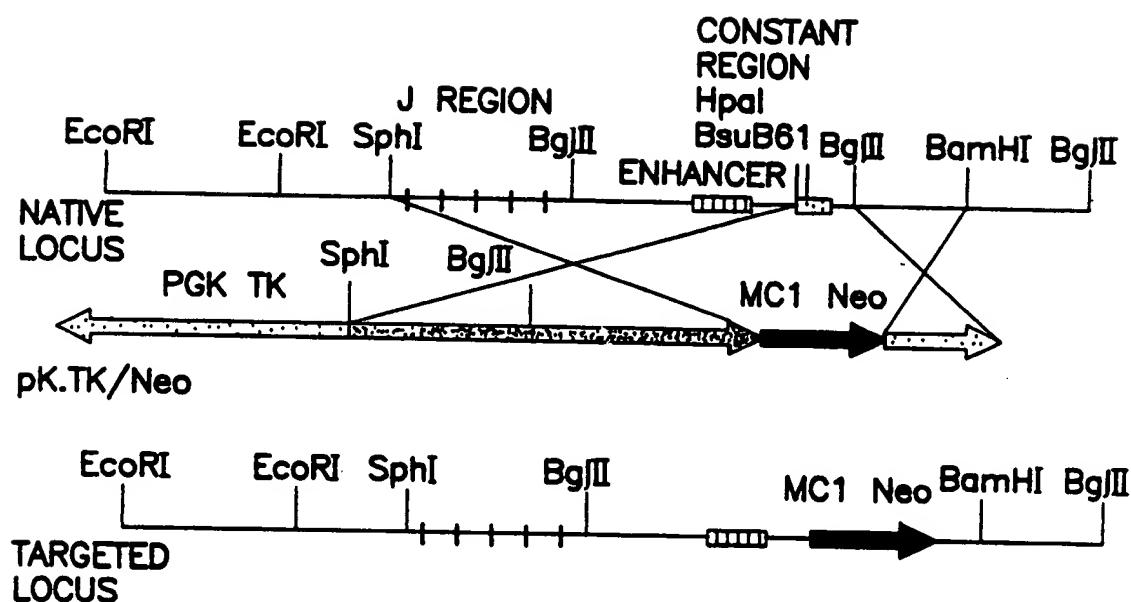
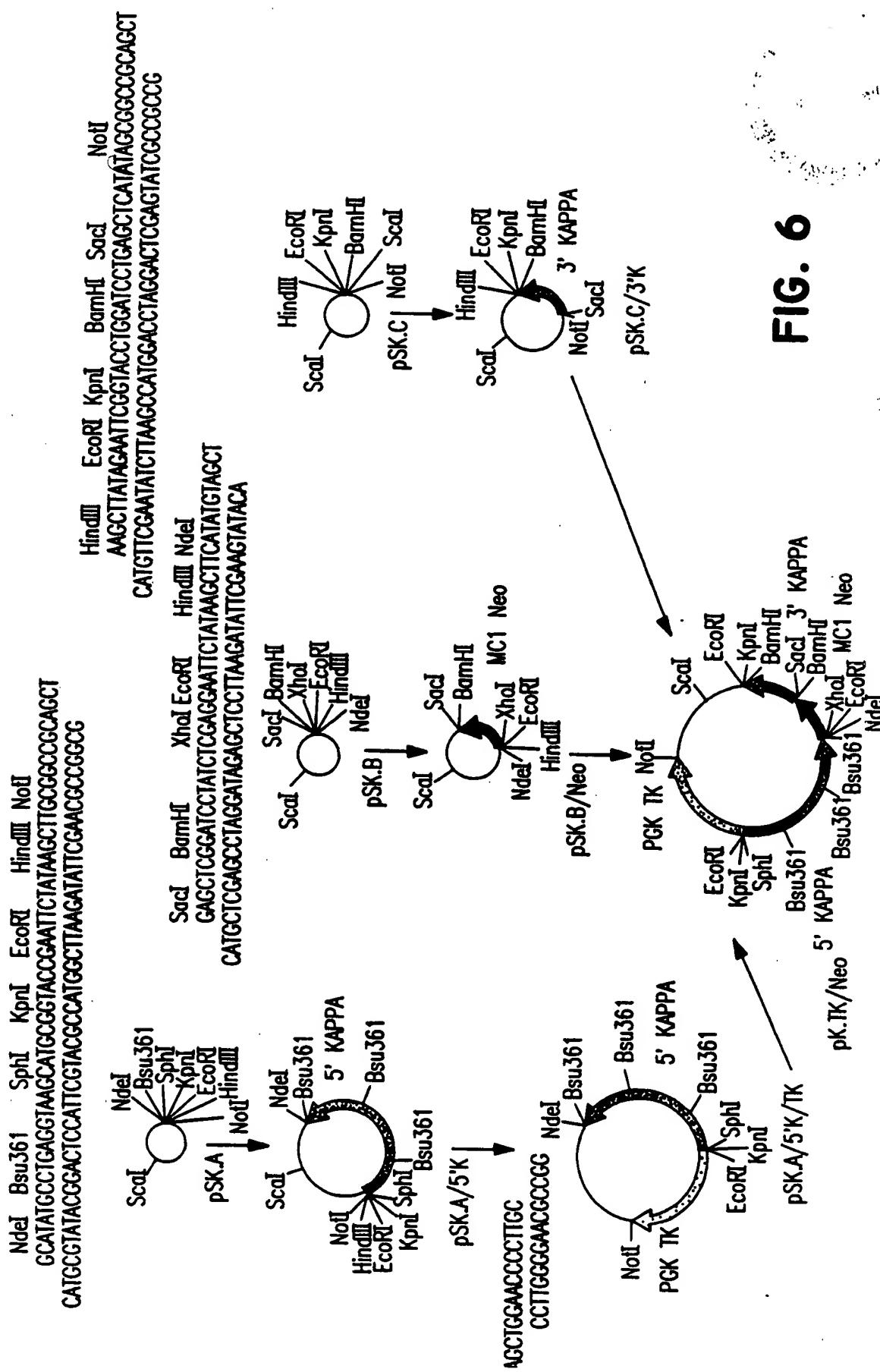
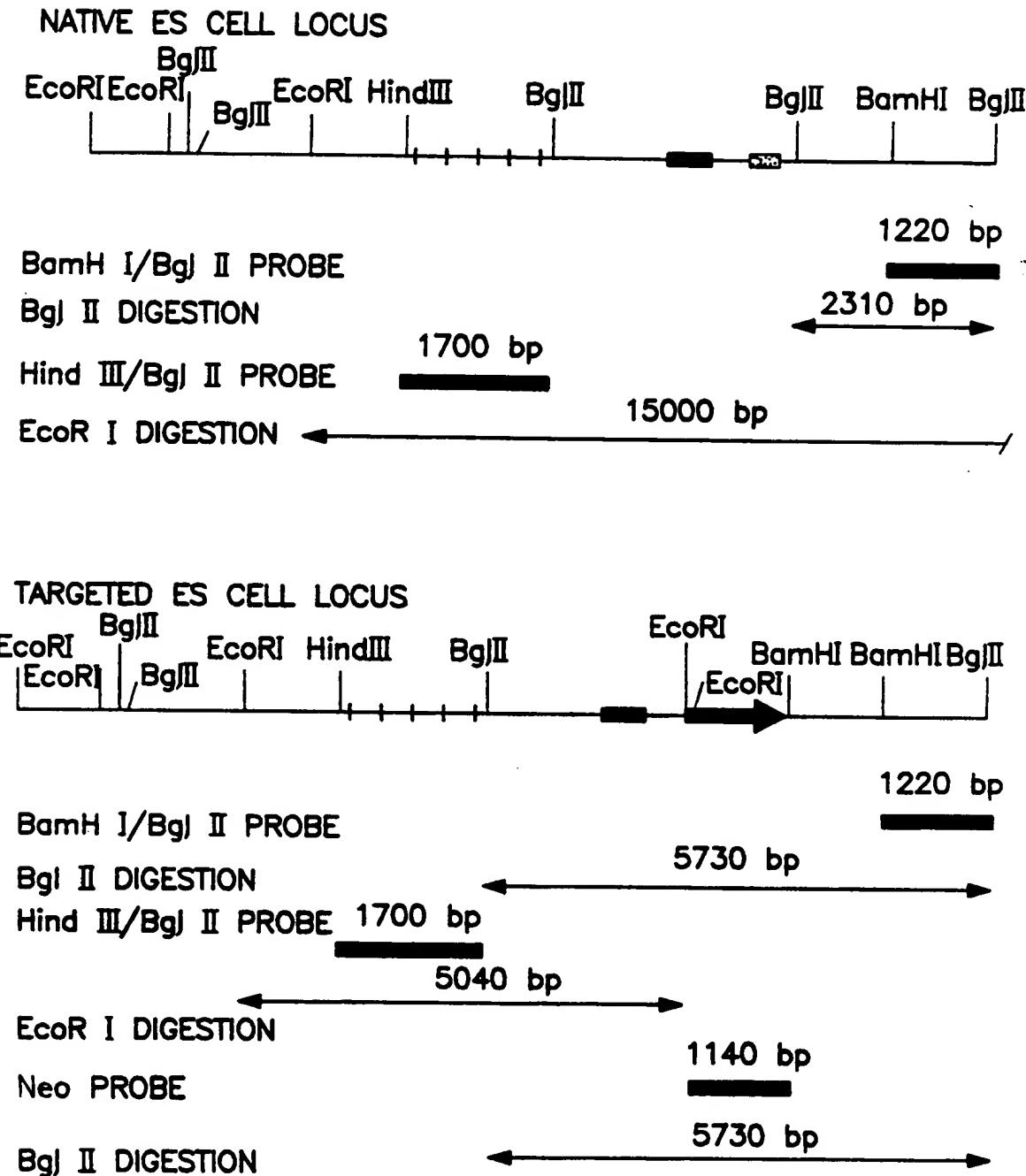
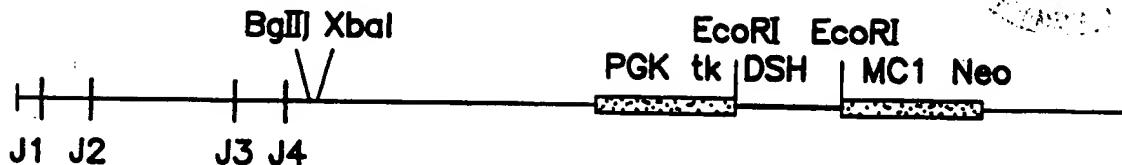


FIG. 5



**FIG. 7**

J REGION KNOCKOUT VECTOR



TARGETING SCHEME

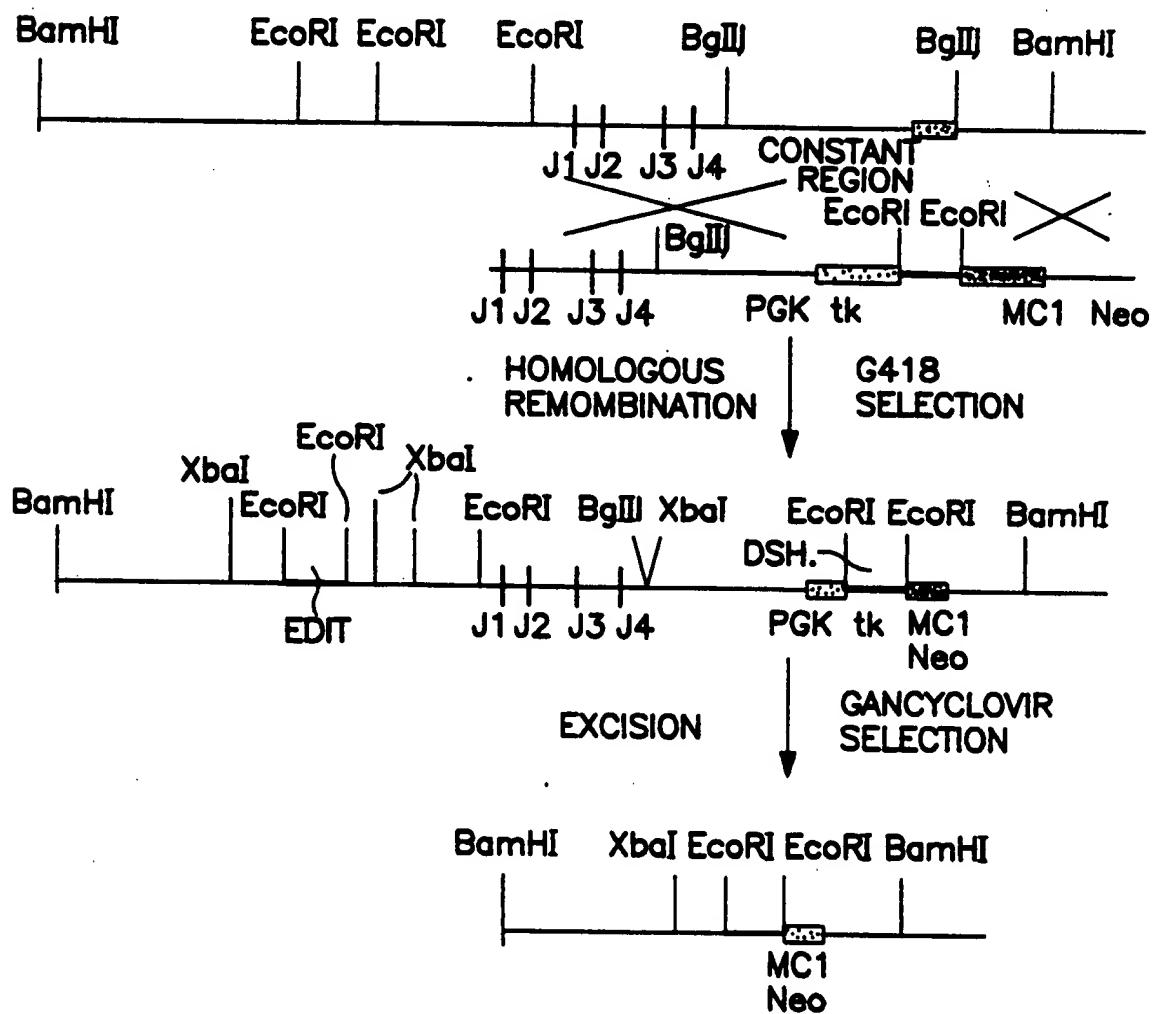


FIG. 8

SacI BamHI XbaI EcoRI HindIII NdeI
 GAGCTCGGATCCTATCTCGAGGAATTCTATAAGCTTCATATGTAGCT
 CATCCTCGAGCCTAGGATAGAGCTCCTTAAGATATTCGAAGTATACA

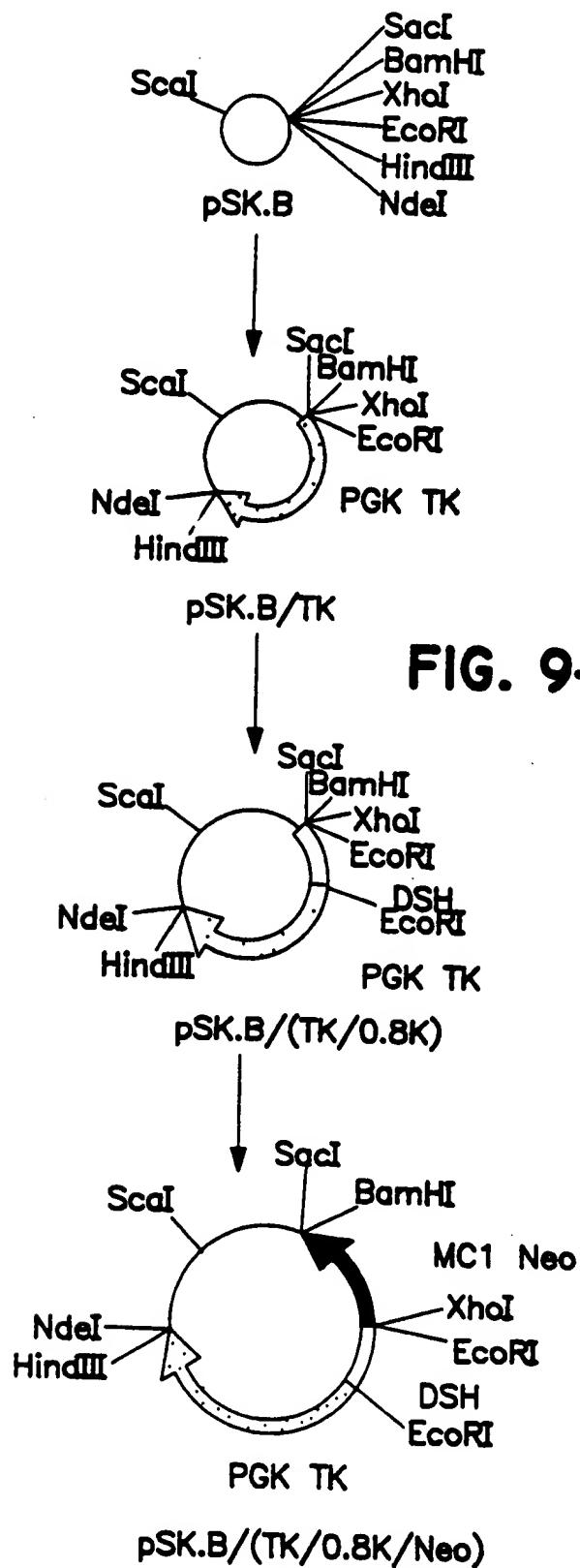


FIG. 9-2

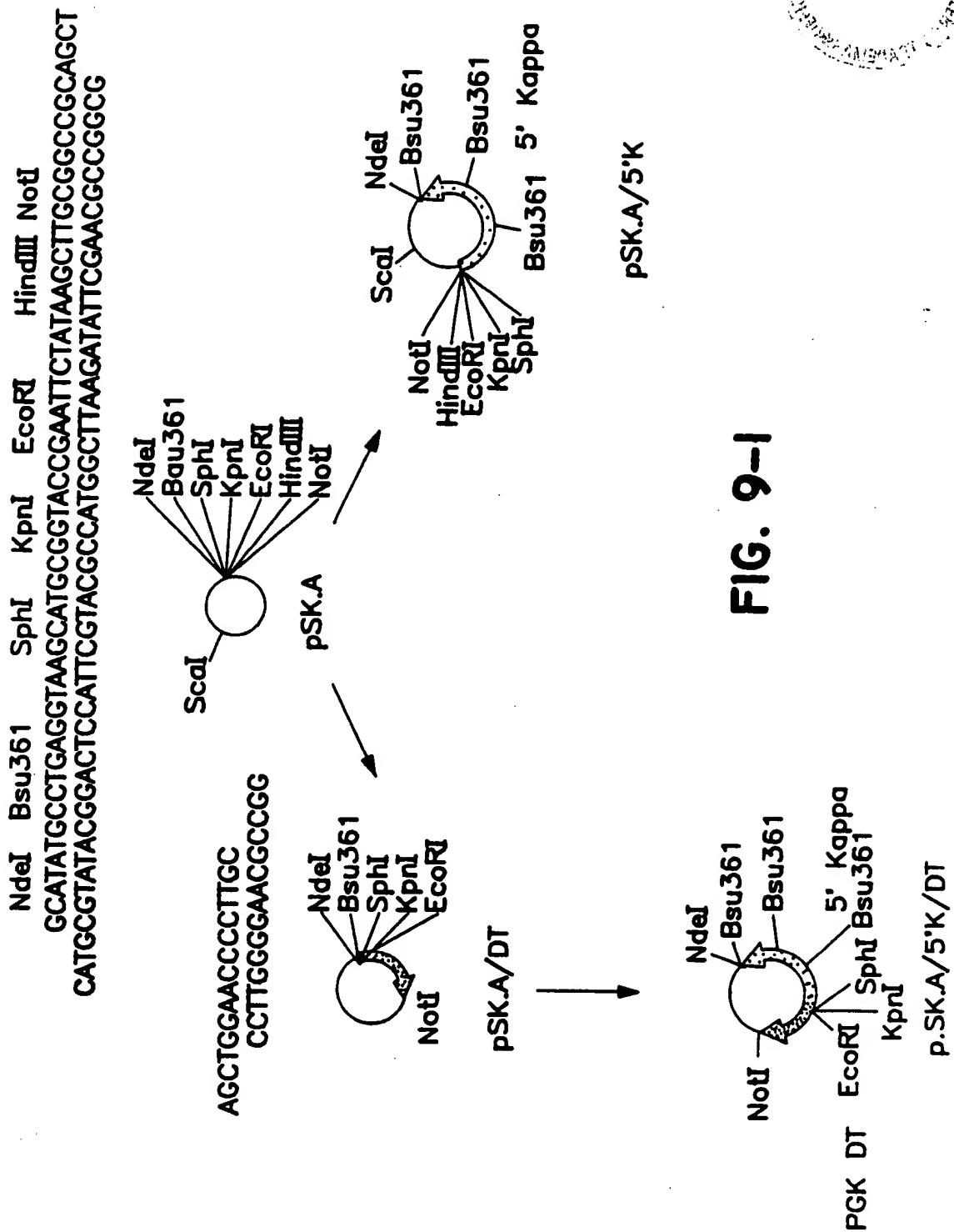


FIG. 9

Hind^{III} EcoRI KpnI BamHI SacI NotI
 AAGCTTATAGAATTGGTACCTGGATCCTGAGCTCATAGCGGCCGCAGCT
 CATGTTGAATATCTTAAGCCATGGACCTAGGACTCGAGTATGCCGGCG

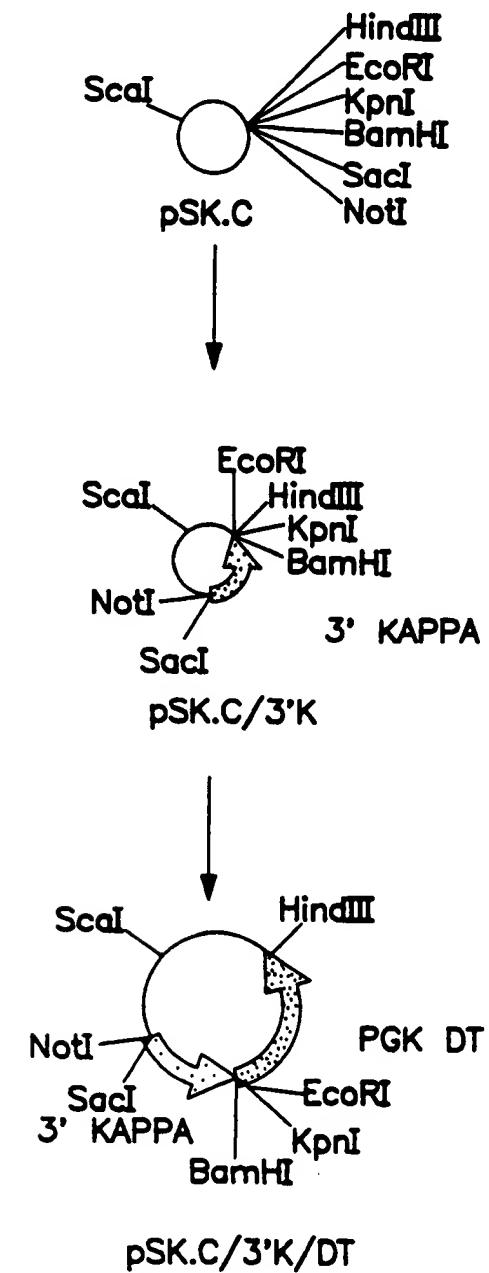
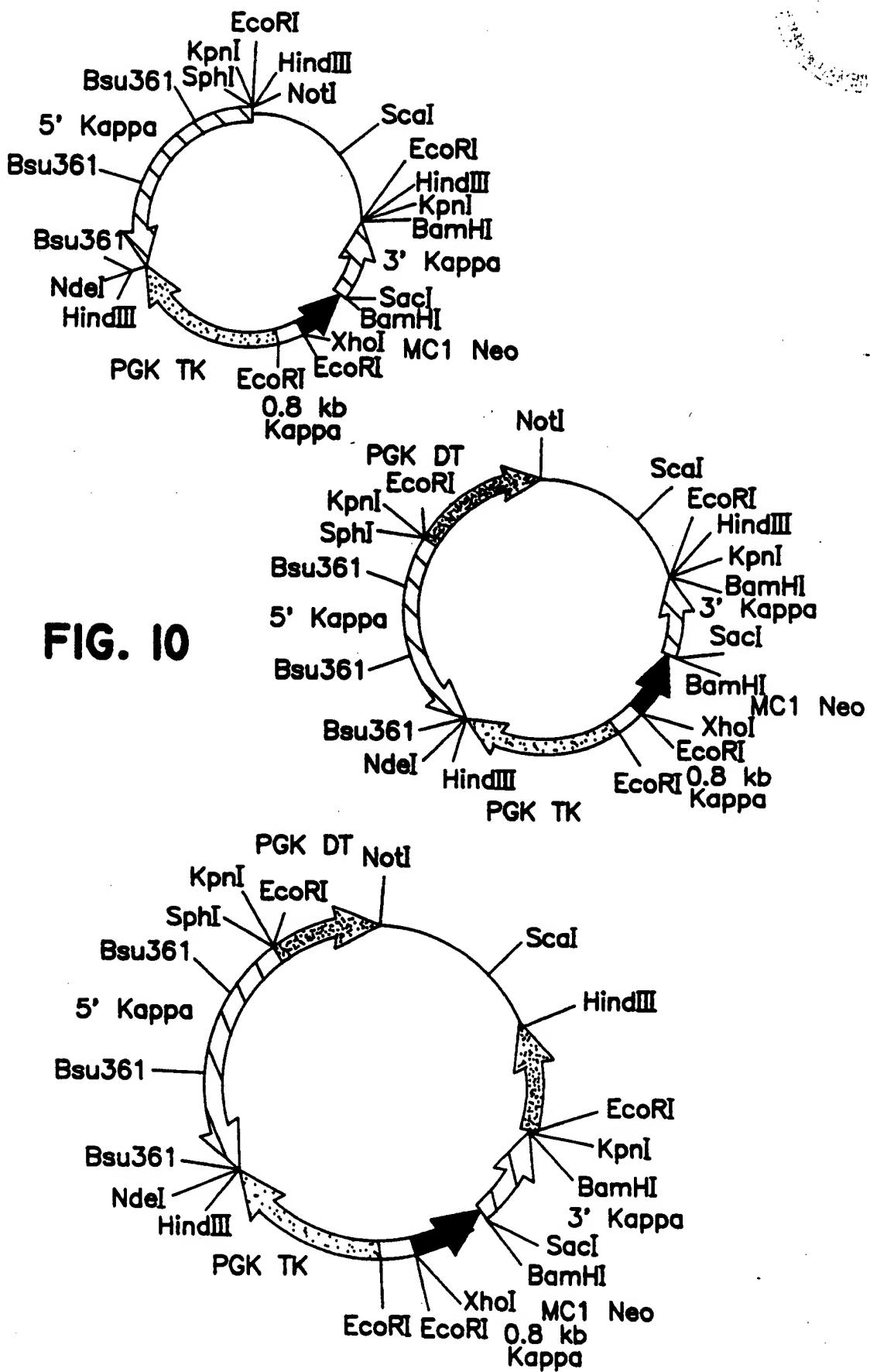
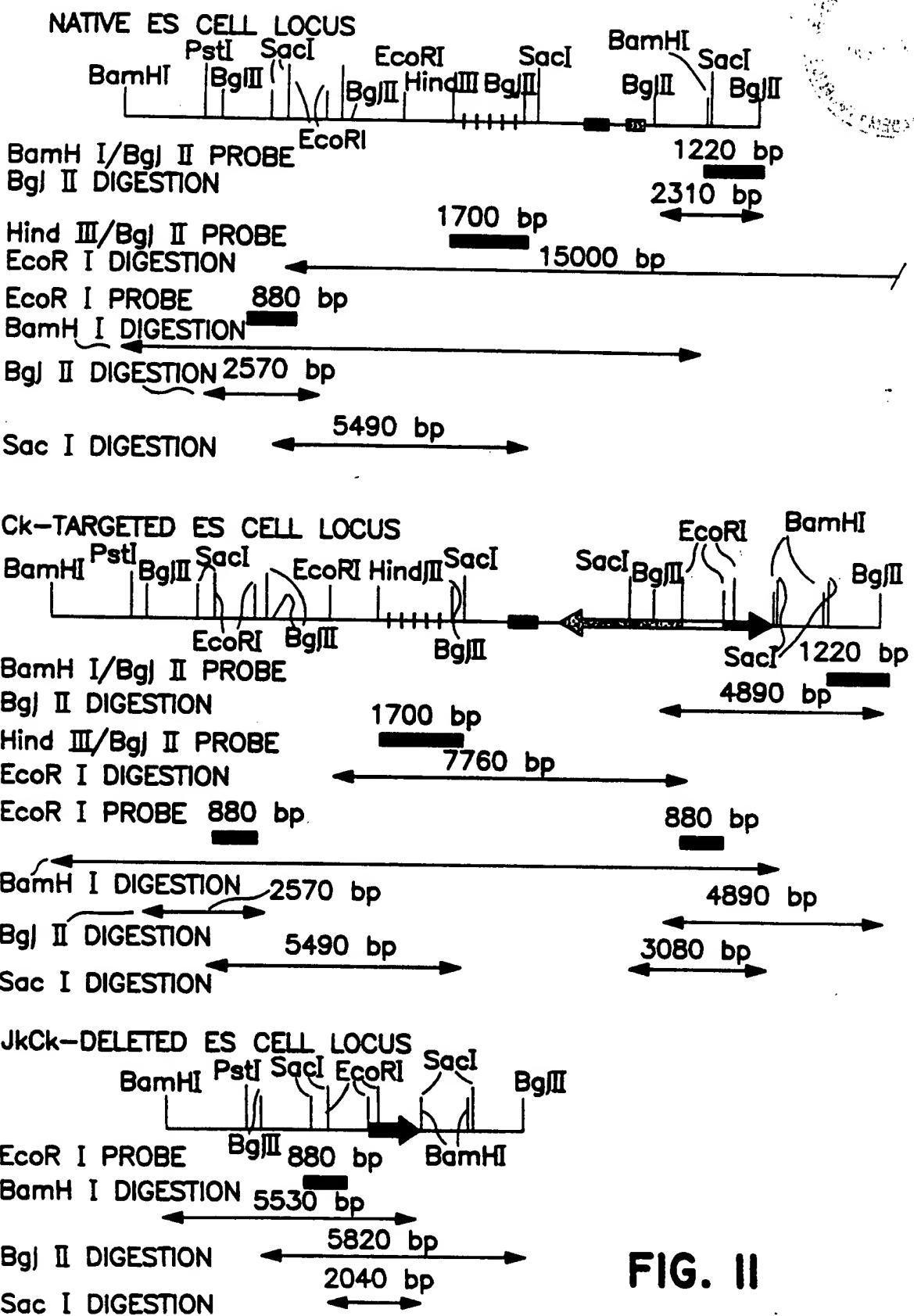


FIG. 9-3



**FIG. II**

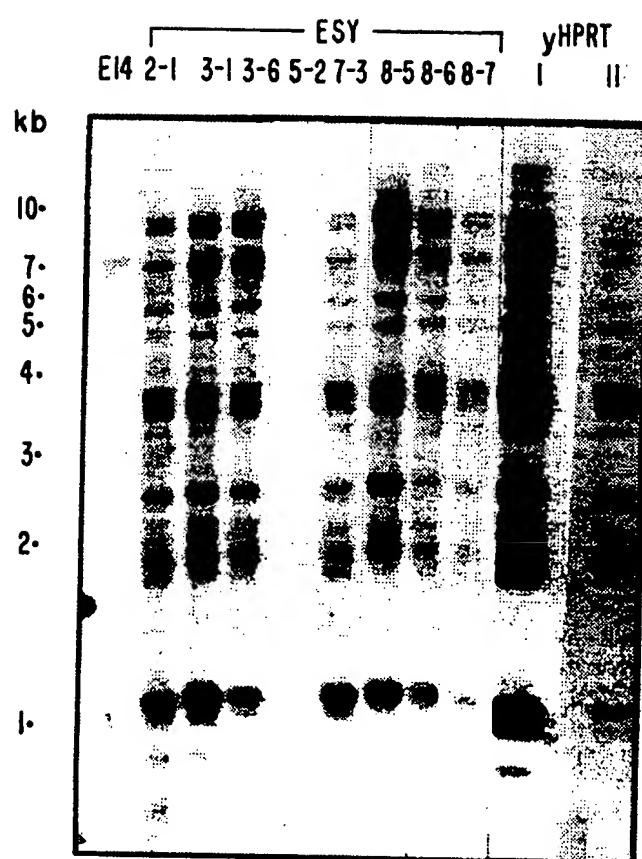


FIG. 12A

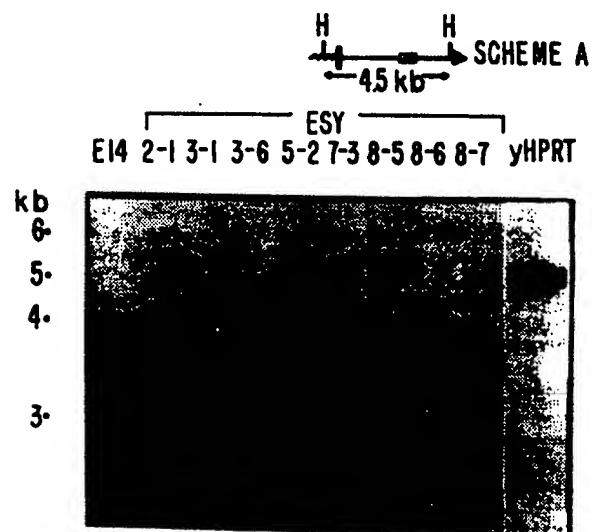


FIG. 12B

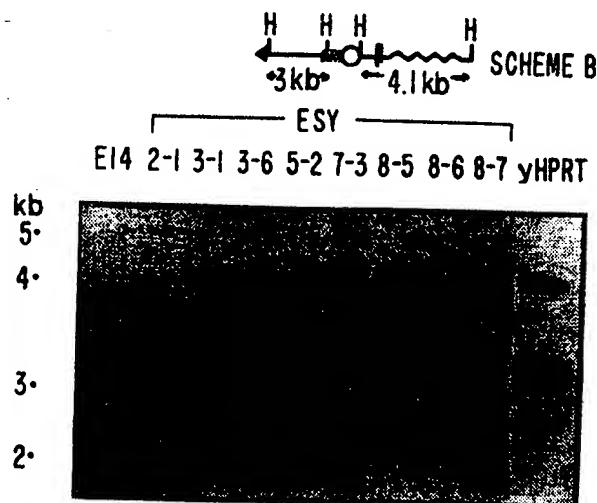
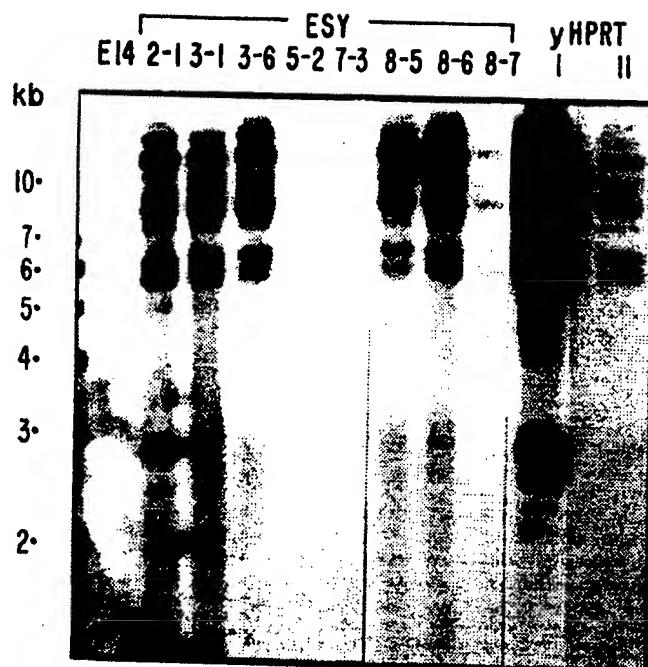
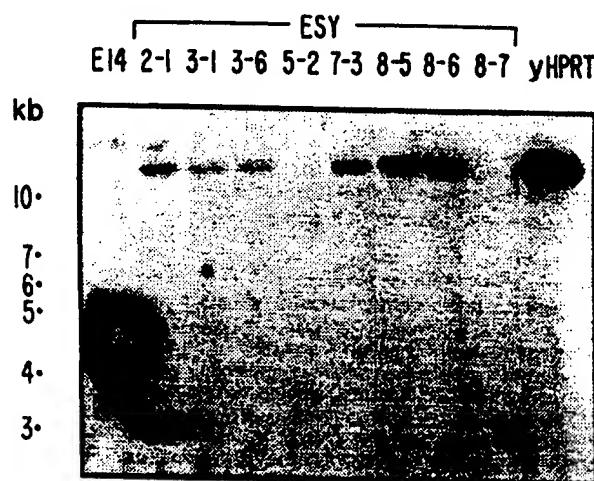
**FIG. 12C****FIG. 12D****FIG. 12E**

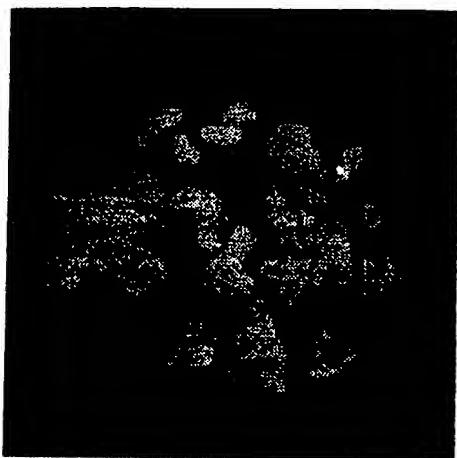
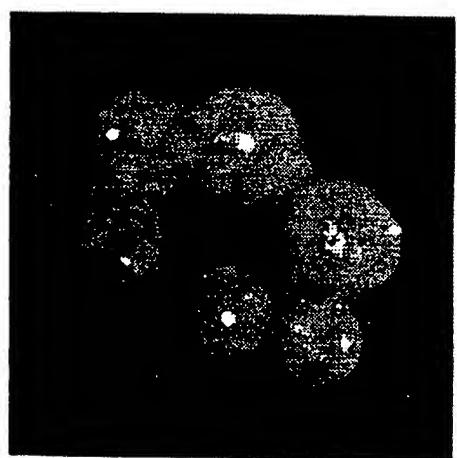
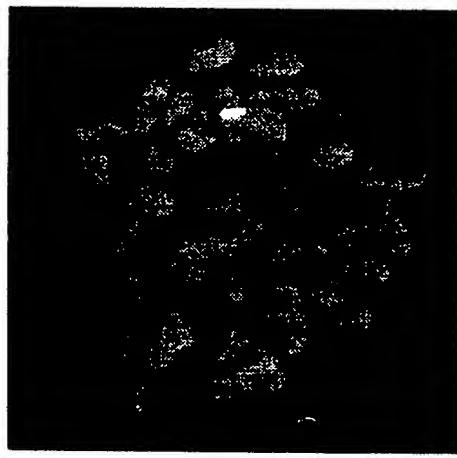
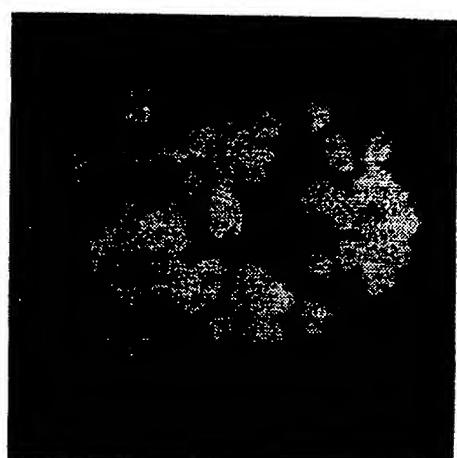
FIG. I3A**FIG. I3B****FIG. I3C****FIG. I3D**



FIG. I4A



FIG. I4B



FIG. I4C



FIG. I4D

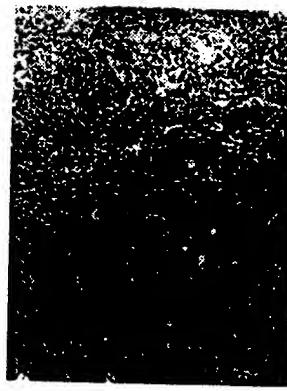


FIG. I4E

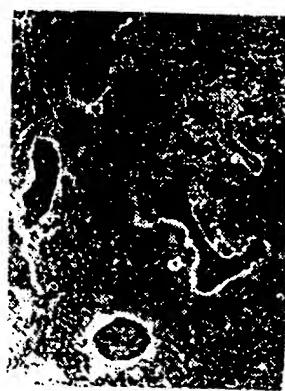


FIG. I4F

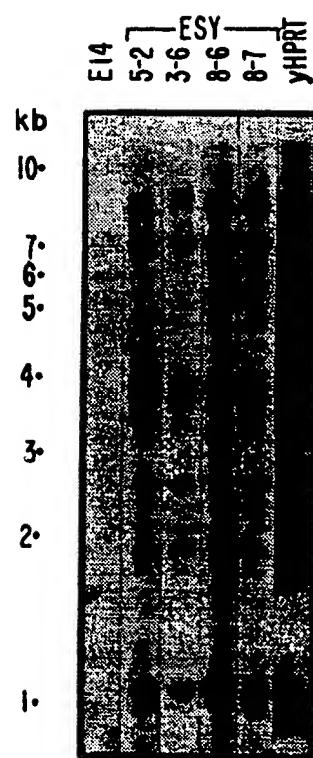


FIG. 14G

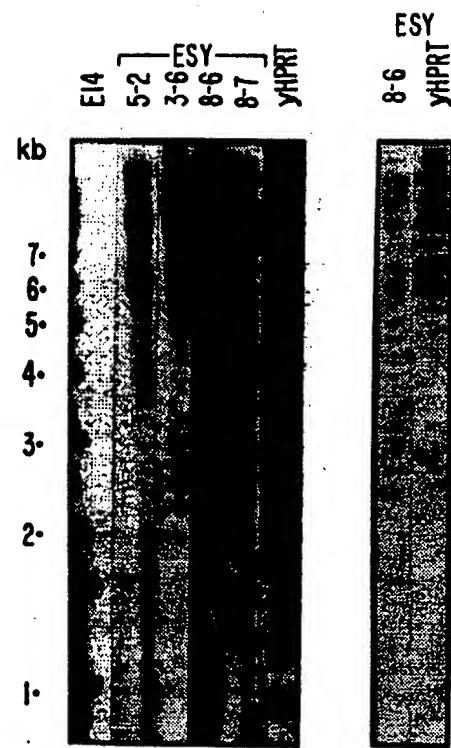
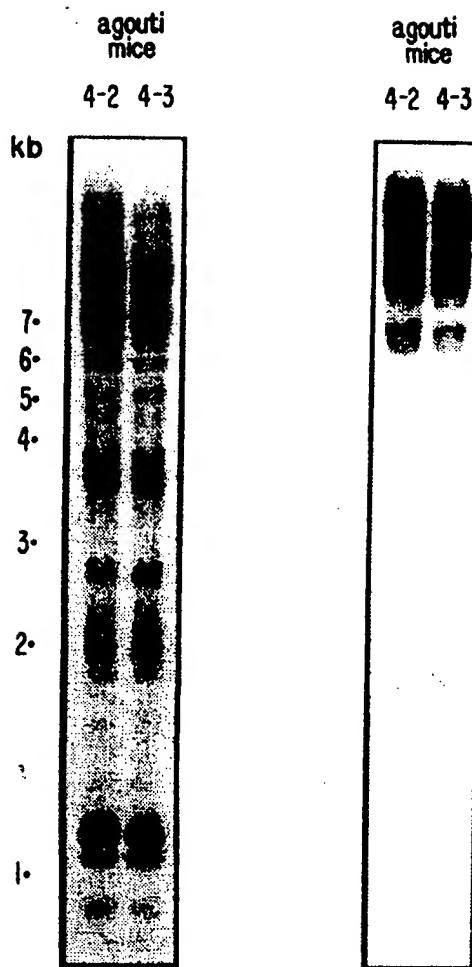
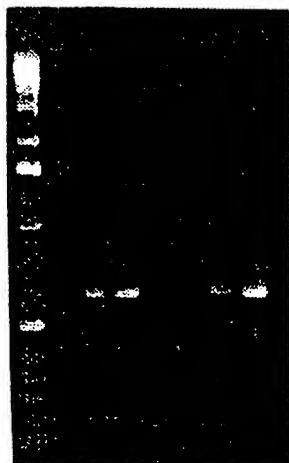


FIG. 14H FIG. 14I

**FIG. 14J****FIG. 14K****FIG. 15A**

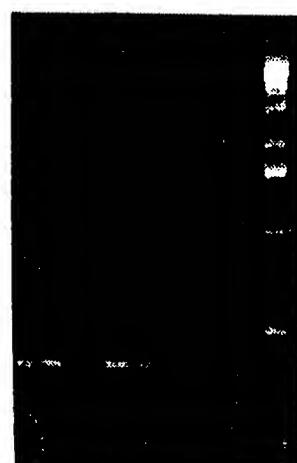
M ES ESY 3-1
Hut 78 C-Liver
C-Spleen 4-3 Liver
4-3 Spleen No DNA



626 bp

FIG. 15B

ES ESY 3-1
Hut 78 C-Liver
C-Spleen 4-3 Liver
4-3 Spleen No DNA
M



359 bp

INTERSPARSED MEMBERS OF V1.V2.V3.V4.V5

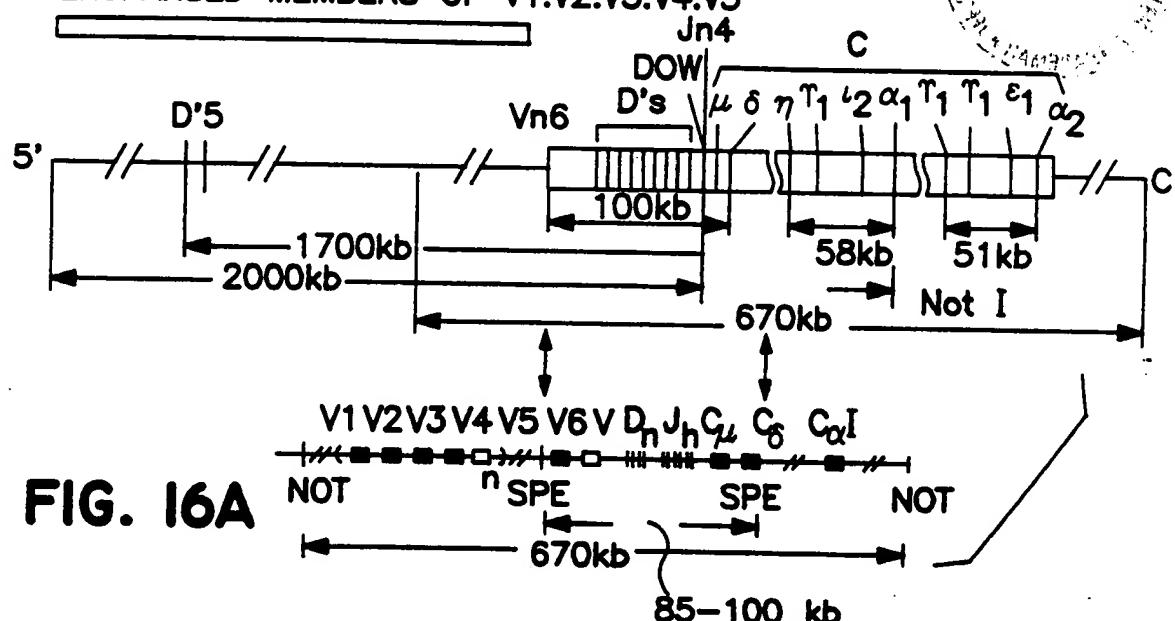


FIG. 16A

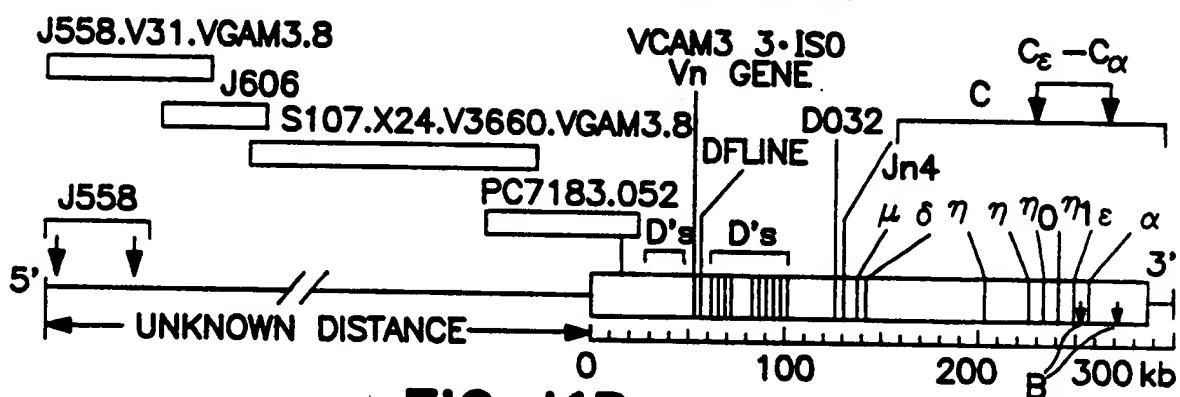


FIG. 16B

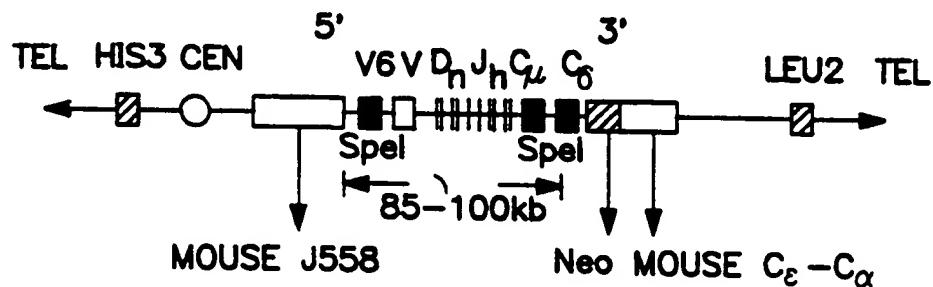


FIG. 16C

سُقُونَاتٍ

Mouse Breeding Scheme

Cross IA.

heterozygous inactive Murine IgH

X

heterozygous inactive Murine IgK

MIGH (inactive) MIGKMIGH MIGK

X

MIGH MIGK (inactive)MIGH MIGK

↓

F1 (cross I A)

MIGH (inactive) MIGK (inactive)MIGH MIGK

Cross II.

F1 (cross I A) x F1 (cross I B)



F2 Quadruple Heterozygotes

MIGH (inactive) MIGK (inactive) HIGH HIGKMIGH MIGK

Cross III.

Intercross F2 mice



F3 DOUBLE Homozygotes

MIGH (inactive) MIGK (inactive) HIGH HIGKMIGH (inactive) MIGK (inactive)

FIG. 17

FIG. 18A

MAMMALIAN HOST GENOTYPES

Intercross Product Mice

- I. Animal I Animal I

II. Animal II Animal II

III. Animal III Animal III

IV. Animal IV Animal IV

V. Animal V Animal V

VI. Animal VI Animal VI

VII. Animal VII Animal VII

VIII. Animal VIII Animal VIII

VIIII. Animal IX Animal IX

X. Animal X Animal X

XI. Animal XI Animal XI

XII. Animal XII Animal XII

XIII. Animal XIII Animal XIII

XIV. Animal XIV Animal XIV

XV. Animal XV Animal XV

XVI. Animal XVI Animal XVI

XVII. Animal XVII Animal XVII

XVIII. Animal XVIII Animal XVIII

XIX. Animal XIX Animal XIX

XX. Animal XX Animal XX

IX.	Animal III	X	Animal IV	$\Delta mIgH$	$mIgH$	$mIgH$	$mIgH$	$mIgH$
X.	Animal II	X	Animal IX	$\Delta mIgH$	$mIgH$	$mIgH$	$mIgH$	$mIgH$
XI.	Animal I	X	Animal IX	$\Delta mIgH$	$mIgH$	$mIgH$	$mIgH$	$mIgH$

*Not all possible genotypes from intercrosses are shown.

Δ = functionally inactive locus
 m = mouse endogenous gene
 h = human transgene
 IgH = immunoglobulin heavy chain
 IgL = immunoglobulin light chain

FIG. 18B